

SEQUENCE LISTING

GENERAL INFORMATION

(i) APPLICANT: Lezdey, John

Lezdey, Darren Kronis, Anne

(ii) TITLE OF INVENTION: DNA FOR EXPRESSION OF ALPHA

1-ANTITRYPSIN IN METHYLOTROPIC

YEAST

(iii) NUMBER OF SEQUENCES: 8

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: John Lezdey & Associates

(B) STREET: 2875 MCI DRIVE (C) CITY: PINELLAS PARK

(D) STATE: FLORIDA (E) COUNTRY: U.S.A. (F) ZIP: 33782

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: CD-R

(B) COMPUTER: Hewlett Packard
(C) OPERATING SYSTEM: WINDOWS 2000
(D) SOFTWARE: Microsoft Word

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 10/672,177 (B) FILING DATE: 9/26/2003

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 09/529,401 (B) FILING DATE: 3/15/2000

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Lezdey, John

(B) REGISTRATION NUMBER: 22,735

(C) REFERENCE/DOCKET NUMBER: 1435-1

(ix)	TELECOMMUNICATION INFORMATION:
(A) (B)	TELEPHONE: (727) 456-2585 TELEFAX: (727) 456-2589
(2)	INFORMATION FOR SEQ ID NO: 1
(i)	SEQUENCE CHARACTERISTICS
(A) (B) (C) (D)	LENGTH: 32 TYPE: NUCLEIC ACID STRANDEDNESS: SINGLE TOPOLOGY: UNKNOWN
(ii)	MOLECULE TYPE: DNA (Genomic DNA)
(iii)	HYPOTHETICAL: No
(iv)	ANTI-SENSE: No
(vi)	ORIGINAL SOURCE
(A)	ORGANISM: HUMAN
(ix)	FEATURE:
	(C) IDENTIFICATION METHOD: gene scan
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 1: CTC GAG AAG AGA GAG GAT CCC CAG GGA GAT GC 32
(2)	INFORMATION FOR SEQ ID NO: 2:
(i)	SEQUENCE CHARACTERISTICS:
(A) (B) (C) (D)	LENGTH: 32 TYPE: NUCLEIC ACID STRANDEDNESS: SINGLE TOPOLOGY: UNKNOWN
(ii)	MOLECULE TYPE: DNA (Genomic DNA)
(iii)	HYPOTHETICAL: No

•	(iv)	ANTI-SENSE: No
:	(vi)	ORIGINAL SOURCE
	(A)	ORGANISM: HUMAN
	(ix)	FEATURE:
		(C) IDENTIFICATION METHOD: gene scan
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 2: GC GGC CGC TTA TTT TTG GGT GGG ATT CAG CAG 32
	(2)	INFORMATION FOR SEQ ID NO: 3
	(i)	SEQUENCE CHARACTERISTICS:
	(A)	LENGTH: 24
	(B)	TYPE: NUCLEIC ACID
		STRANDEDNESS: SINGLE
	(D)	TOPOLOGY: UNKOWN
	(ii)	MOLECULE TYPE: DNA (Genomic DNA)
	(iii)	HYPOTHETICAL: No
	(iv)	ANTI-SENSE: No
	(vi)	ORIGINAL SOURCE
	(C)	ORGANISM: HUMAN
	(ix)	FEATURE:
	(D)	IDENTIFICATION METHOD: gene scan
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 3: CAG AAG ACA GAT ACA TCC CAC CAT 24
	(2)	INFORMATION FOR SEQ ID NO: 4:
	(i)	SEQUENCE CHARACTERISTICS:
	(4)	LENGTU. 22
	(A)	LENGTH: 22
	(B)	TYPE: NUCLEIC ACID
	(C)	STRANDEDNESS: SINGLE

(D)	TOPOLOGY: UNKNOWN	
(ii)	MOLECULE TYPE: DNA (Genomic DNA)	
(iii)	HYPOTHETICAL: No	
(iv)	ANTI-SENSE: No	
(vi)	ORIGINAL SOURCE	
(A)	ORGANISM: HUMAN	
(ix)	FEATURE:	
(B)	IDENTIFICATION METHOD: gene scan	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 4: AGG ATT TCA TCG TGA GTG TCA G	22
(2)	INFORMATION FOR SEQ ID NO: 5:	
(i)	SEQUENCE CHARACTERISTICS:	
(A) (B) (C) (D)	LENGTH: 24 TYPE: NUCLEIC ACID STRANDEDNESS: SINGLE TOPOLOGY: UNKNOWN	
(ii)	MOLECULE TYPE: DNA (Genomic DNA)	
(iii)	HYPOTHETICAL: No	
(iv)	ANTI-SENSE: No	
(vi)	ORIGINAL SOURCE	
(A)	ORGANISM: HUMAN	
(ix)	FEATURE:	
(B)	IDENTIFICATION METHOD: gene scan	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 5:	24

:	(2)	INFORMATION FOR SEQ ID NO: 6:
	(i)	SEQUENCE CHARACTERISTICS:
	(B) (C)	LENGTH: 24 TYPE: NUCLEIC ACID STRANDEDNESS: SINGLE TOPOLOGY: UNKNOWN
	(ii)	MOLECULE TYPE: DNA (Genomic DNA)
	(iii)	HYPOTHETICAL: No
	(iv)	ANTI-SENSE: No
	(vi)	ORIGINAL SOURCE
	(A)	ORGANISM: HUMAN
	(ix)	FEATURE:
	(B)	IDENTIFICATION METHOD: gene scan
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 6: AGC TTC TTA CAG TGC TGG ATG TTA 24
	(2)	INFORMATION FOR SEQ ID NO: 7:
	(i)	SEQUENCE CHARACTERISTICS:
	(A) (B) (C) (D)	LENGTH: 24 TYPE: NUCLEIC ACID STRANDEDNESS: SINGLE TOPOLOGY: UNKNOWN
	(ii)	MOLECULE TYPE: DNA (Genomic DNA)
	(iii)	HYPOTHETICAL: No
	(iv)	ANTI-SENSE: No
	(vi)	ORIGINAL SOURCE

•	(A)	ORGANISM: HUMAN	
-	(ix)	FEATURE:	
	(B)	IDENTIFICATION METHOD: gene scan	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 7: GTT CAA CAA ACC CTT TGT CTT CTT	24
	2)	INFORMATION FOR SEQ ID NO: 8:	
	(i)	SEQUENCE CHARACTERISTICS:	
	(A) (B) (C) (D)		
	(ii)	MOLECULE TYPE: DNA (Genomic DNA)	
	(iii)	HYPOTHETICAL: No	
	(iv)	ANTI-SENSE: No	
	(vi)	ORIGINAL SOURCE	
	(A)	ORGANISM: HUMAN	
	(ix)	FEATURE:	
	(B)	IDENTIFICATION METHOD: gene scan	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 8: GGG AGA CTT GGT ATT TTG TTC AAT	24